SEQUENCE LISTING

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Trifillis, Panayiota

<120> METHODS FOR IDENTIFYING COMPOUNDS THAT MODULATE UNTRANSLATED REGION-DEPENDENT GENE EXPRESSION AND METHODS OF USING SAME

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<141> 2004-01-21 (371c date)

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<151> 2004-01-21

<150> 60/441,637
<151> 2003-01-21

<160> 90

<170> PatentIn version 3.2

<210> 1
<211> 14
<212> DNA
<213> Artificial Sequence

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<223> Description of Artificial Sequence: consensus G-quartet element from synthetic sequences

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<221> misc_feature
<222> 3, 7, 8, 11
<223> n = a, t, c, or g

<220>
<221> misc_feature
<222> (7)..(8)
<223> This represents one form of the sequence as described, other forms described may have up to five nucleotides in this variable region

<400> 1
ggntggnngg ntgg 14

<210> 2
<211> 14
<212> DNA
<213> Artificial Sequence

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<223> Description of Artificial Sequence: synthetic G-quartet oligonucleotide

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<222> 3, 4, 7, 8, 11, 12
<223> n = a, t, g or c

<220>
<221> misc_feature
<222> 3, 4, 7, 8, 11, 12
<223> This represents one form of the sequence as described, other forms described have longer variable regions, typical is 2 - 10 nucleotides

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<210> 3
<211> 61
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Antisense minus uORF NcoI primer

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g 61

<210> 4
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<223> subunit of 15-LOX-DICE

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ccccrcccuc uucccccaag 19

<210> 5
<211> 152
<212> DNA
<213> Homo sapiens

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ggctccaccc tctctccccc ggaaaggaca cc 152

<210> 6
<211> 792
<212> DNA
<213> Homo sapiens

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<210> 7
<211> 21
<212> RNA
<213> Homo sapiens

<220>
<223> Group I AU-Rich element(ARE) cluster of 3'untranslated region

<400> 7
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<210> 8
<211> 40
<212> DNA
<213> Homo sapiens

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<210> 9
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<212> DNA
<213> Homo sapiens

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accctgatac aggcatggca gaagaatggg aatattttat actgacagaa atcagtaata 180
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<212> DNA
<213> Artificial Sequence

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<212> DNA
<213> Homo sapiens

<400> 11
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<210> 12
<211> 307

<212> DNA
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<223> Group III AU-Rich element(ARE) cluster of 3'untranslated region

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<210> 14
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<212> DNA
<213> Homo sapiens

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<220>
<223> Description of Artificial Sequence: Expression Vector pCMR2

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<212> DNA
<213> Homo sapiens

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<212> DNA
<213> Homo sapiens

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<211> 1153
<212> DNA
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<211> 334
<212> DNA
<213> Homo sapiens

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<210> 36
<211> 543
<212> DNA
<213> Homo sapiens

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<210> 37
<211> 511
<212> DNA
<213> Homo sapiens

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<210> 38
<211> 458
<212> DNA
<213> Homo sapiens

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<210> 39
<211> 270
<212> DNA
<213> Homo sapiens

<400> 39
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<210> 40
<211> 751
<212> DNA
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<221> misc_feature
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<223> n = a, t, g or c

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<210> 41
<211> 229
<212> DNA
<213> Homo sapiens

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<210> 42
<211> 233
<212> DNA
<213> Homo sapiens

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<210> 43

<211> 349
<212> DNA
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<210> 44
<211> 337
<212> DNA
<213> Homo sapiens

<400> 44
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<212> DNA
<213> Homo sapiens

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<210> 69
<211> 35
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer 1 for amplify VEGF 5'UTR

<400> 69
aaagtcgacg taatcgcgga ggcttgggc agccgg 35

<210> 70
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer 2 for amplify VEGF 5'UTR

<400> 70
tttgcgactg gtcagctgcg ggatcccaag 30

<210> 71
<211> 33
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer 3 for amplify VEGF 5'UTR

<400> 71
aagtcgacgt aagagctcca gagagaagtc gag 33

<210> 72
<211> 33
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer 4 for amplify VEGF 5'UTR

<400> 72
aaacccgggc agcaaggcaa ggctccaatg cac 33

<210> 73
<211> 39
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer 5 for amplify VEGF 3'UTR

<400> 73
gccgggcagg aggaaggagc ctccctcagg gtttgggga 39

<210> 74
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer 6 for amplify VEGF 3'UTR

<400> 74
ctgcactaga gacaaagacg tgatgttaat 30

<210> 75
<211> 66
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Polylinker

<400> 75
gaacaaatgt cgacgggggc ccctaggaga tctagcgctg gatcccccgg ggagctcaug 60

gaagac 66

<210> 76
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer for luciferase amplification

<400> 76
cggtgttggg cgcgttattt atcggagttg 30

<210> 77
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer for luciferase amplification

<400> 77
ttggcgaaga atgaaaatag ggttggtact 30

<210> 78

<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer for GAPDH amplification

<400> 78
ggtgaaggtc ggagtcaacg ga 22

<210> 79
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer for GAPDH amplification

<400> 79
gagggatctc gctcctggaa g 21

<210> 80
<211> 55
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: 5'UTR forward oligo

<400> 80
aaagtcgacg taaccgccag atttgaatcg cgggacccgt tggcagaggt ggcgg 55

<210> 81
<211> 54
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: 5'UTR reverse oligo

<400> 81
aaaggatccg ggcaacgtcg gggcacccat gccgccgccg ccacctctgc caac 54

<210> 82
<211> 40
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: 3'UTR forward oligo

<400> 82
aaagcggccg cggcctctgc cggagctgcc tggtcccaga 40

<210> 83
<211> 37

<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: 3'UTR reverse oligo

<400> 83
aaatctagac tcaggaacag ccgagatgac ctccaga 37

<210> 84
<211> 67
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: SL top oligonucleotide

<400> 84
ctagaagctt agggccgcgg atccgcgcgc ggttcgccgc gcgcggatcc gcggtagcaa 60
gttagtc 67

<210> 85
<211> 68
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: SL bottom oligonucleotide

<400> 85
gactaagctt gctaccgcgg atccgcgcgc ggcgaaccgc gcgcggatcc gcggccctaa 60
gcttctag 68

<210> 86
<211> 32
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR primer (Sense/HindIII)

<400> 86
caagaagctt gcgcccggcc ccccaccccc cg 32

<210> 87
<211> 31
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR primer (Antisense/NcoI)

<400> 87
agcccatggt gctcactgcg gctccggccc c 31

<210> 88

<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR primer (Sense/BglII)

<400> 88
agactctgaa ccagaaggcc aa 22

<210> 89
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR primer (Antisense/KpnI)

<400> 89
ctcggtacca gttttccaaa atatatttgc aaatgg 36

<210> 90
<211> 58
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: sense minus uORF HindIII primer

<400> 90
cccaagcttc gcgcccggcc ccccaccccc cgcagcaccc cgcgccccgc gccctccc 58